

SEQUENCE LISTING

<110> Ben-Sasson, Shmuel

<120> Tissue Remodeling

<130> K-08-US

<160> 59

<170> PatentIn version 3.1

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Gln Thr Leu

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<211> 10
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<400> 24

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<400> 32

Glu Ile Ala Arg Arg Cys Ser Ile Gly Gly Ile His Glu Asp Tyr Gln
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Leu Pro Tyr Tyr Asp Leu Val Pro Ser Asp Pro Ser Val Glu Glu
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<210> 33
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 <212> PRT
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<400> 33

Glu Leu Val Ser Arg Cys Lys Ala Ala Asp Gly Pro Val Asp Glu Tyr
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Met Leu Pro Phe Glu Glu Glu Ile Gly Gln His Pro Ser Leu Glu Glu
 20 25 30

<210> 34
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<213> unknown-synthetic

<400> 34

Glu Ile Phe Met Arg Cys Thr Asp Leu Phe Pro Gly Glu Ser Val Pro
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Glu Tyr Gln Met Ala Phe Gln Thr Glu Val Gly Asn His Pro Thr Phe
20 25 30

Glu Asp

<210> 35

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<212> PRT

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<400> 35

Glu Met Thr Ser Arg Cys Asn Ala Val Gly Glu Val Lys Asp Tyr Glu
1 5 10 15

Pro Pro Phe Gly Ser Lys Val Arg Glu His Pro Cys Val Glu Ser
20 25 30

<210> 36

<211> 31

<212> PRT

<213> unknown-synthetic

<400> 36

Glu Ile Ala Arg Arg Cys Asn Ser Gly Gly Val His Glu Glu Tyr Gln
1 5 10 15

Leu Pro Tyr Tyr Asp Leu Val Pro Ser Asp Pro Ser Ile Glu Glu
20 25 30

<210> 37

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<400> 37

Glu Met Ala Arg Arg Cys Ile Thr Gly Gly Ile Val Glu Glu Tyr Gln
1 5 10 15

Leu Pro Tyr Tyr Asn Met Val Pro Ser Asp Pro Ser Tyr Glu Asp
 20 25 30

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<400> 38

Glu Ile Ala Arg Arg Thr Ile Val Asn Gly Ile Val Glu Asp Tyr Arg
 1 5 10 15

Pro Pro Phe Tyr Asp Val Val Pro Asn Asp Pro Ser Phe Glu Asp
 20 25 30

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<400> 39

Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
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Thr Val Thr Val Glu
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<400> 40

Thr Ala Phe His Asp Lys Gly Ser Leu Thr Asp Tyr Leu Lys Gly Asn
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Ile Ile Thr Trp Asn Glu
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<400> 41

Met Glu Tyr Tyr Pro Asn Gly Ser Leu Cys Lys Tyr Leu Ser Leu His
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Thr Ser Asp Trp Val
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<210> 42
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<400> 42

Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg His
 1 5 10 15

Val Ile Ser Trp Glu
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<210> 43
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<400> 43

Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
 1 5 10 15

Thr Val Thr Ile Glu
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<210> 44
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<400> 44

Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Phe Leu Lys Cys Ala
 1 5 10 15

Thr Leu Asp Thr Arg
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<400> 45

Thr	His	Tyr	His	Glu	His	Gly	Ser	Leu	Tyr	Asp	Phe	Leu	Gln	Arg	Gln
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Thr	Leu	Glu	Pro	His
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<400> 46

Ala	Asp	Asn	Lys	Asp	Asn	Gly	Thr	Trp	Thr	Gln	Leu	Trp	Leu	Val	Ser
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Asp

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Ala	Glu	Lys	Arg	Gly	Ser	Ser	Leu	Glu	Ala	Glu	Leu	Trp	Leu	Ile	Thr
1				5					10					15	

Ala

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Gly	Asp	Glu	Arg	Val	Thr	Ala	Asp	Gly	Arg	Met	Glu	Tyr	Leu	Leu	Val
1				5					10					15	

Met Glu

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 1 5 10 15

Ala

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Ala Asp Asn Lys Asp Asn Gly Thr Trp Thr Gln Leu Trp Leu Val Ser
 1 5 10 15

Asp

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Ala Asp Ile Lys Gly Thr Gly Ser Trp Thr Gln Leu Tyr Leu Ile Thr
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Asp

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<400> 52

Ser Asp Met Thr Ser Arg Asn Ser Ser Thr Gln Leu Trp Leu Ile Thr
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His

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Gln Thr Val Met Leu Arg His Glu Asn Ile Leu Gly Phe Ile Ala
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Ser Thr Pro Gly Met Lys His Glu Asn Leu Leu Gln Phe Ile Ala
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Arg Val Pro Leu Met Glu His Asp Asn Ile Ala Arg Phe Ile Val
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Ser Asp Ile Asn Leu Lys His Glu Asn Ile Leu Gln Phe Leu Thr
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<400> 57

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